

6

No.	Target	File	Definition	Match%	Over.	INIT	OPT
2	SmtW_H_DNA_N			82.8	372	198	1066
			10 20 30 40 50 60				
SmtW_R_ORF.dn	ATGCTTGCCAGGGGCCGAGCGGC	CCCCCGCCGCGCCCGCCTCCGGTCTTTCCCTTC	X:::	:	:	:	:
SmtH_H_DNA_N	ATGCCGGCCTCTGCTGCGCGGCCCGCCGCGGTCCC	GCGACGCTACAGCCTCGCCCTTC					
	10 20 30 40 50 60						
	70 80 90 100 110 120						
SmtW_R_ORF.dn	CCGCGCGCCGCTGCTGCTGCTGCTGCTGCTGCGATACTAAGCGCGCCGCTGTCGCGCGC	::	:	:	:	:	:
SmtH_H_DNA_N	CC-----GCTACTGCTGCTGCGCGTCTGAGCGCGCCGCTATCCGGCGCGC						
		70 80 90 100					
	130 140 150 160 170 180						
SmtW_R_ORF.dn	GTCCCCCGCTCAGTGCGCGAGAACCCTGCTGCGCATCTCCGAGGCTGA	CCTATCTCAGC	:	:	:	:	:
SmtH_H_DNA_N	GTCCCCCGCTCGGTGCGCGAGAACCCTCGCTTCCAATCTCTGAGGCTGA	CCTCTGCTCAGC					
	110 120 130 140 150 160						
	190 200 210 220 230 240						
SmtW_R_ORF.dn	CGGTTTTGACGCGTCTCATACGTACAATTACTGTGCTCTCCTTGTG	GAATCCTGCCTCCAC	:	:	:	:	:
SmtH_H_DNA_N	CGGTTCCGAGTCCCTCACACATACAATTACTGTGTTCTCCTTGTG	AATCCTGCCTCCAC					
	170 180 190 200 210 220						
	250 260 270 280 290 300						
SmtW_R_ORF.dn	ACACITTAACGTCGGTGACGGGATAGCATGTCGCTTTAACCTCCCCTTCTG	TGGGGAA	:	:	:	:	:
SmtH_H_DNA_N	ACACITTAATGTTGGCGCCGGGACACCATCTTCGCTTTATCCCTGCGCTTCTG	AGGGAG					
	230 240 250 260 270 280						
	310 320 330 340 350 360						
SmtW_R_ORF.dn	AGACCCCGAAGGATCGACTGATGGTAGCTGAGACTCACAGACAGAACTG	CAGGAAGAAA	:	:	:	:	:
SmtH_H_DNA_N	AGACCCCGCAGGATTGACTGGATGGTTCTGAGGCTCACAGACAGAACTGT	AGGAAGAAA					
	290 300 310 320 330 340						
	370						
SmtW_R_ORF.dn	GGCAAGAAAGAG						
SmtH_H_DNA_N	GGCAAGAAAGAG						